



## SEQUENCE LISTING

&lt;110&gt; Institut National de la Recherche Agronomique (INRA)

&lt;120&gt; Method of producing double low restorer lines of Brassica napus having a good agronomic value

&lt;130&gt; D21413

&lt;140&gt; US 10/563,277

&lt;141&gt; 2006-01-04

&lt;160&gt; 14

&lt;170&gt; PatentIn version 3.2

&lt;210&gt; 1

&lt;211&gt; 248

&lt;212&gt; DNA

&lt;213&gt; Brassica napus

&lt;220&gt;

&lt;223&gt; PGIol marker

&lt;400&gt; 1

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acatgtgggt	aacttaacag	ggctccggct	gttgcaaaac	acatgggtgc	tgctcagcact	120
aatcttgcgg	tatgaatttg	tgattaaaatt	tgtttggttg	tgactctttc	ttcattgttc	180
gttttcgtac	aataaacgca	atgtataatc	ttttacaaa	ctgaattttc	taccgggtct	240
gatgtaca						248

&lt;210&gt; 2

&lt;211&gt; 979

&lt;212&gt; DNA

&lt;213&gt; Brassica napus

&lt;220&gt;

&lt;223&gt; PGI-UNT R2000 marker

&lt;400&gt; 2

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gtaagtgcct	gtttatttgg	ttgtataaat	ttctcgtcca	tttccgcttg	cttagtgtat	480
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agttgtgtga	ttatacagtt	ttcttgcctt	tttgcctatg	ccatcaacac	tagagattcg	900
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 <223> PGI-int R2000 marker

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 ggggtctgatg tacaatgcta gtctccatgt tcttggggat catgatttat tttctacatg 180  
 tattcagaca gtacagaaga aagtgttcaa aactctggat gttttaattt acagtttagt 240  
 gagaagttcg gcattgatcc gaacaatgca ttgtcatatt gggactgggt tgggtggaagg 300  
 tacagtggta agtgcttgtt tattttggtg tataaatttc tctgccattt ccgcttgcctt 360  
 agtgataaac tgaaattctt ttgcagtttg cagtgtctgt ggagtcttac cattgtctct 420  
 acagattggc ttctctgttg ttgagaagta cggtaacctc tactttatca gccatctcat 480  
 aaaaatgtctt aggcataatc ttcttatttt atttccctct taatgatttc ttcttttttt 540  
 tattgcattc ccgttttatt ttcaaaagtt gttactgtct ctaaatcaag aagaaacctt 600  
 cttagtagat ccagctgata ttcagccttt tttaaattgg actgcaggtt tttaaagggg 660  
 agcttcaagc attgataagc atttccagtc cacaccgttt gagaagaata taccctgtag 720  
 ttgtcattgt tgtgtgatta tacagttttc ttgtcttttt gctatgtcca tcaacactag 780  
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 <212> DNA  
 <213> Brassica napus

<220>  
 <223> BolJon marker R2000

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 gggataacag tgtagaaaac aaaccgtctg taagattttc tccctgatcc tctcaacttaa 180  
 ccagtagggc tttttcacat tgaagcgcat atctactttg gtatctactg aataaaaaaa 240  
 gaaagctggg aacatgtgaa ggatatacaa gcattgatac accaagtagt cacaactac 300  
 attataaagg tcagaccttt gttcacattc tggcctccag gaccaccgct tctagcaaa 360  
 ttaagcgtaa catgggtctgc acgtatacaa atgaaaatgt tttctatcaa atctctataa 420  
 atagagctct ataacattgt cgatacatag ttccactaac tctgcaagta ctaaacacat 480  
 atcaaaaaca aactatgcga acagatcaaa actactacag aacacagttc tatgacactg 540  
 tcgatatgta catctctctg aagtacccaa gagatagcaa atgaaactat gtaaaaact 600  
 caaaatctca aatttctcca tcacaaggac ctacagaata gagttatcat aacattttct 660  
 gtaaatattt catcaaaaat gactagagaa cagagtctct ataacattat ctgtaaatgt 720  
 tccaaacaaa ccactacata cgagagttct tataacattg tctgtaaatg tccaatcaaa 780  
 accactacga acaaaaagctc ctataacatt gtttatcaaa agtttcaact aatctacaaa 840  
 ctttccccgt aatgagctt aatatcacc aaagatgttt caatcagata aagagtacga 900  
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 <223> CP418L marker R2000

<400> 5

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ccactacata gcagagttct tataacattg tctgtaaatg tccaatcaaa accactacag 180
aacaagaagctc ctataacatt gtttatacaa agtttcaact aatctacaaa ctttccccgt 240
aaatgagctt aatatcacc aaagatgttt caatcagata aagagtaacg acatcggttt 300
gagattagaa caaaactgaaa cttacgtaga gtgatttgag gagtaggctc gttgccagca 360
gagctagctc tctctctcgc ctcacgaagc atctgttgca cctgagacaa ccgtgacgaa 420
actttccgat caccgccacc agaattcgac gccgcgcac ggaaggatcc gaatcgggaa 480
ctgagtgaa cccgagcgatc ccgggagtg gcaggagcga tgggaaaaa gagtgggcacg 540
atttcgacga agagtggag agggaggggt ggtggataaa ctgcgcgtat atcaagtctc 600
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<213> artificial sequence

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<220>
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<400> 6
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<220>
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<220>
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<400> 9
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 <212> DNA  
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 <400> 11  
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